

Figure 1A
Prostate Derived Ets Factor

1	GTCTGACTTCCTCCCAGCACATTCTCTGCACTCTGCGGTGTCCCACTGCCCCACAGACCC	60
61	AGTCCTCCAAGCCTGCTGCCAGCTCCCTGCAAGCCCTCAGGTGGGCTTGCCACGGTG	120
121	CCAGCAGGCAGCCCTGGGCTGGGGGTAGGGGACTCCCTACAGGCACGCAGCCCTGAGACC	180
181	TCAGAGGGCCACCCCTTGAGGGTGGCCAGGCCCTCAGTGGCCAACTGAGTGTCTGCCTCT	240
241	GCCACCAGCCCTGCTGCCCCCTGCTTCCTGCTGGCCCCCAGATGCCTGGCTGAGACACGC	300
301	CAGTGGCCTCAGCTGCCACACCTCTTCCCGGCCCTGAAGTTGGCACTGCAGCAGACAG	360
361	CTCCCTGGGCACCAGGCAGCTAACAGACACAGCCGCCAGCCCAACAGCAGCGGCATGGG	420
1		M G 2
421	CAGCGCCAGCCCGGGTCTGAGCAGCGTATCCCCCAGCCACCTCCTGCTGCCCCCGACAC	480
3	S A S P G L S S V S P S H L L L P P D T	22
481	GGTGTGCGCGACAGGCTTGGAGAAAGCGGCAGCGGGGGTAGTGGGTCTCGAGAGACGGGA	540
23	V S R T G L E K A A A G A V G L E R R D	42
541	CTGGAGTCCCAGTCCACCCGCCACGCCCGAGCAGGGCCTGTCCGCCTTCTACCTCTCCTA	600
43	W S P S P P A T P E Q G L S A F Y L S Y	62
601	CTTTGACATGCTGTACCCTGAGGACAGCAGCTGGGCAGCCAAGGCCCTTGGGGCCAGCAG	660
63	F D M L Y P E D S S W A A K A P G A S S	82
661	TCGGAGGAGCCACCTGAGGAGCCTGAGCAGTGGCCGGTCAATTGACAGCCAAGCCTCAGC	720
83	R E E P P E E P E Q C P V I D S Q A P A	102
721	GGGCAGCCTGGACTTGGTGCCCGCGGGCTGACCTTGGAGGAGCACTCGCTGGAGCAGGT	780
103	G S L D L V P G G L T L E E H S L E Q V	122
781	GCACTCCATGGTGGTGGGCGAAGTGCTCAAGGACATCGAGACGGCCTGCAAGCTGTCAA	840
123	Q S M V V G E V L K D I E T A C K L L N	142
841	CATCACCGCAGATCCCATGGACTGGAGCCCCAGCAATGTGCAGAAGTGGCTCCTGTGGAC	900
143	I T A D P M D W S P S N V O K W L L W T	162

Figure 1B
Prostate Derived Ets Factor

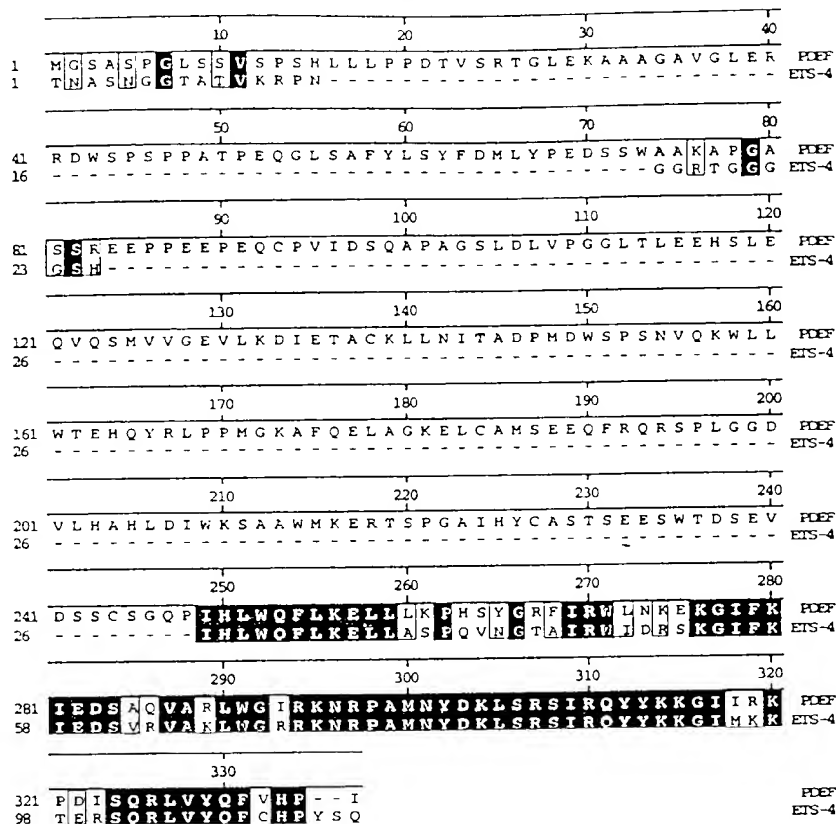
901	AAGACCAATACCGGCTGCCCCCATGGGCAAGGCTTCAGGAGCTGGCGGGCAAGGA	960
163	<u>E H Q Y R L P P M G K A F Q E L A G K E</u>	182
961	GCTGTGCGCCATGTGCGAGGAGCAGTTCGGCCAGCGCTCGCCCTGGGTGGGGATGTGCT	1020
183	<u>L C A M S E E Q F R O R S P L G G D V L</u>	202
1021	GCACGCCACCTGGACATCTGGAAGTCAGCGGCTTGATGAAAAGAGCGGACTTCACCTGG	1080
203	<u>H A H L D I W K S A A W M K E R T S P G</u>	222
1081	GCGATTCACTACTGTGCCTCGACCAGTGAGGAGAGCTGGACCGACAGCGAGGTGGACTC	1140
223	<u>A I H Y C A S T S E E S W T D S E V D S</u>	242
1141	ATCATGCTCCGCGCAGCCCATCCACCTGTGGCAGTTCCTCAAGGAGTTGCTACTCAAGCC	1200
243	<u>S C S G Q P I H L W Q F L K E L L L K P</u>	262
1201	CCACAGCTATG3CCGCTTCATTAGGTG3CTCAACAAGGAGAAG3GCATCTTCAAAATTGA	1260
263	<u>H S Y G R F I R W L N K E K G I F K I E</u>	282
1261	GGACTCAGCCCAGGTGGCCCGGCTGTG3GGCATCCGCAAGAA3CGTCCCGCCATGAACTA	1320
283	<u>D S A Q V A R L W G I R K N R P A M N Y</u>	302
1321	CGACAAGCTGAGCCGCTCCATCCGCCAGTATTACAAGAAGGGATCATCCGGAAGCCAGA	1380
303	<u>D K L S R S I R Q Y Y K K G I I R K P D</u>	322
1381	CATCTCCCAGCGCTCGTCTACCAAGTTCGTGCACCCCATCTGAGTGCCTGGCCAGGGCC	1440
323	<u>I S Q R L V Y Q F V H P I *</u>	336
1441	TGAAACCCGCCCTCAGGGGCTCTCTCCTGCCTGCCTGCCTCAGCCAGGCCCTGAGATG	1500
1501	GGGGAACCGGCGAGTCTGCTCTGCTGCTTGACCTTCAGAGCCCAAGGTGAGGGAGGG	1560
1561	GCAACCAACTGCCCCAGGGGGATATGGGTCTCTGCGGCTTCGGGACCATGGGGCAGGG	1620
1621	GTGCTTCCTCCTCAGGCCCAGCTGCTCCTCGAGGACAGAGGGAGACAGGGCTGCTCCC	1680
1681	CAACACCTGCCTCTGACCCAGCATTTCCAGAGCAGAGCCTACAGAAGGGCAGTGACTCG	1740
1741	ACAAAGGCCACAGGCAGTCCAGGCCTCTCTGCTCCATCCCCCTGCCTCCCATTTCTGCA	1800

Figure 1C
Prostate Derived Ets Factor

1801 CCACACCTGGCATGGTGCAGGGAGACATCTGCACCCCTGAGTTGGGCAGCCAGGAGTGCC 1860

1861 CCCGGAATGGATAATAAGATACTAGAGAACTG 1894

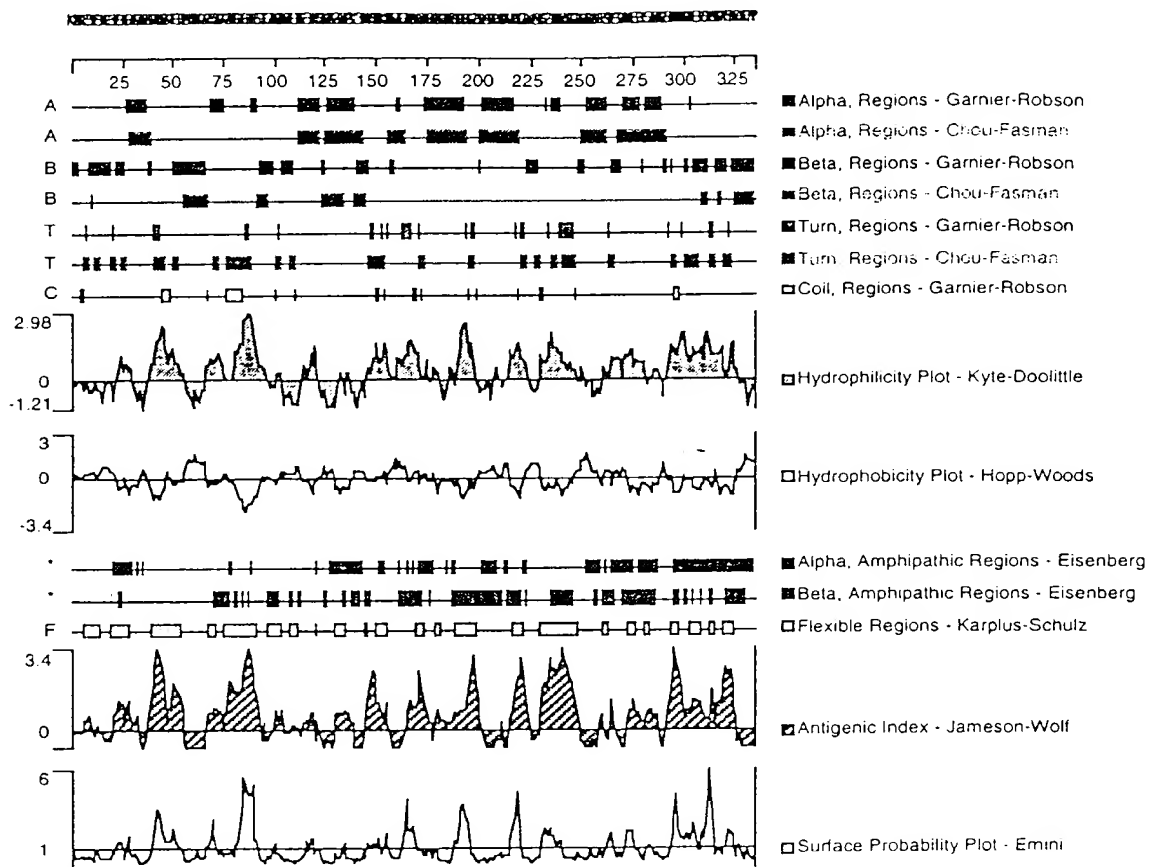
Figure 2



Decoration 'Decoration #1': Box residues that match the consensus named 'Consensus #2' exactly.

Decoration 'Decoration #2': Shade (with solid black) residues that match the consensus named 'Consensus #1' exactly.

Figure 3



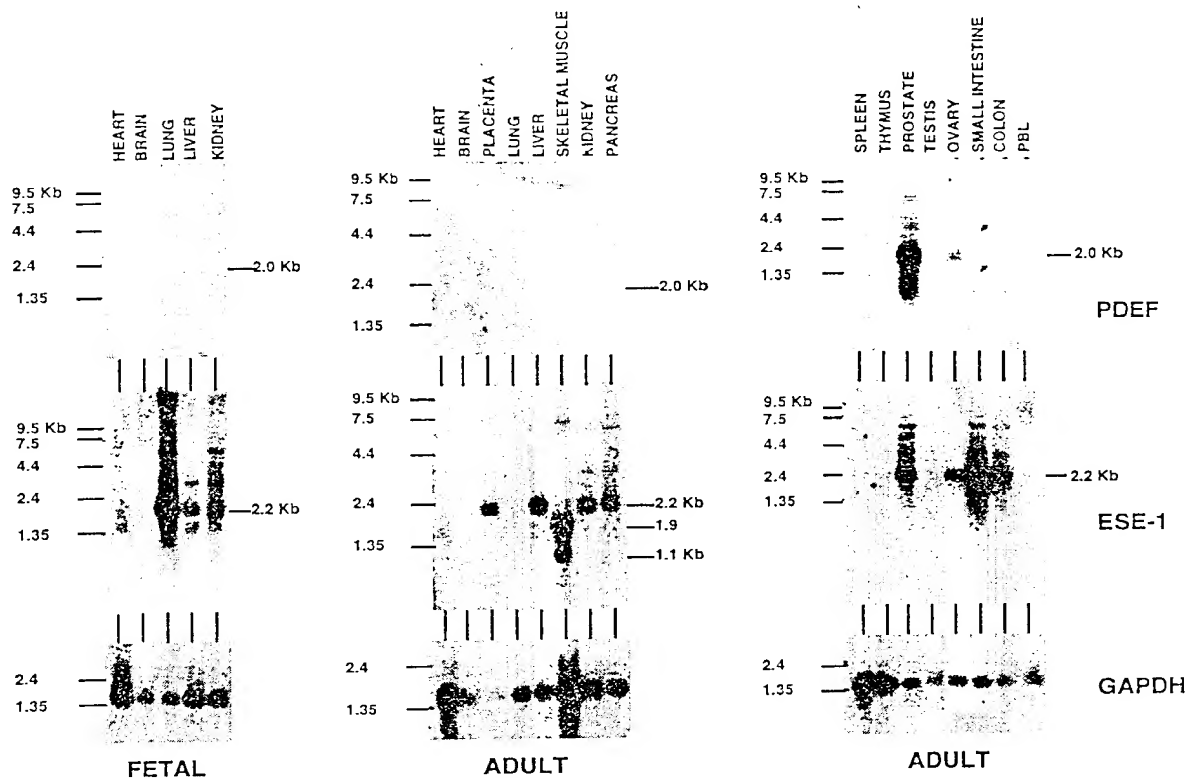


FIGURE 4

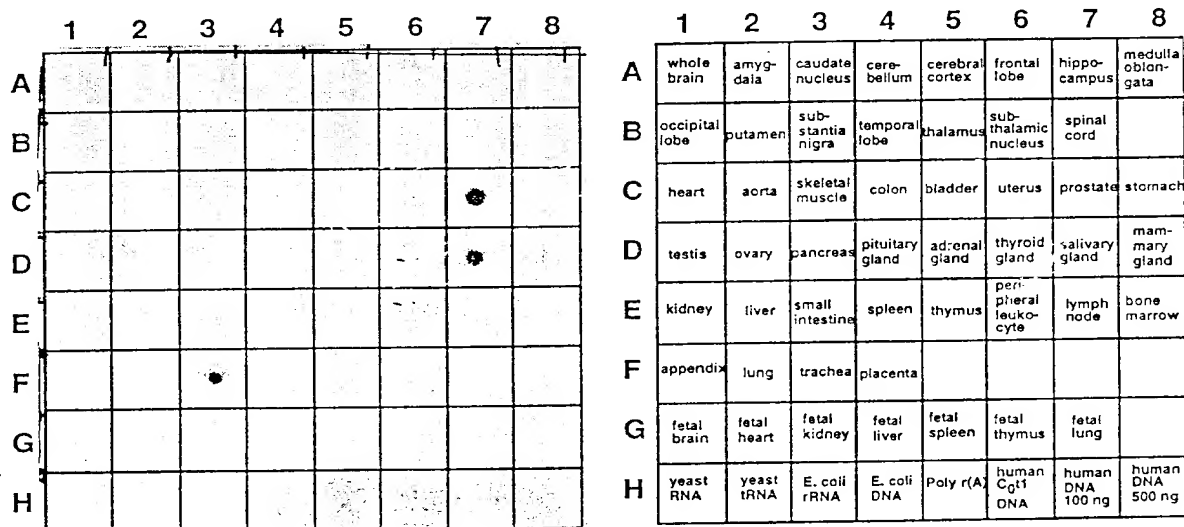


FIGURE 5

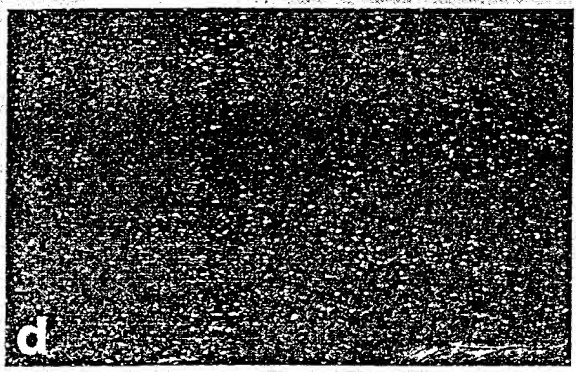
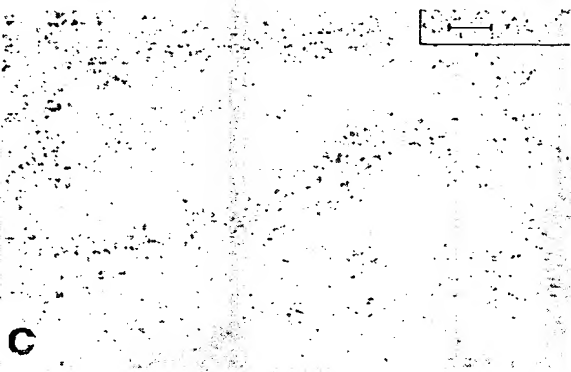
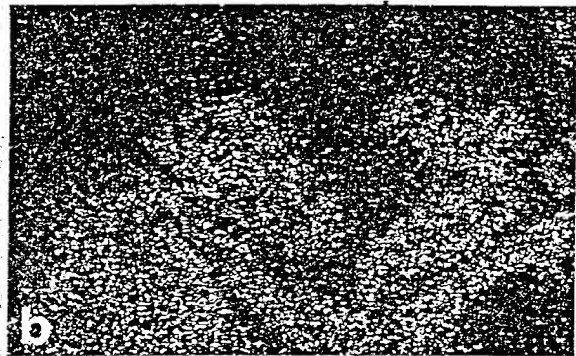
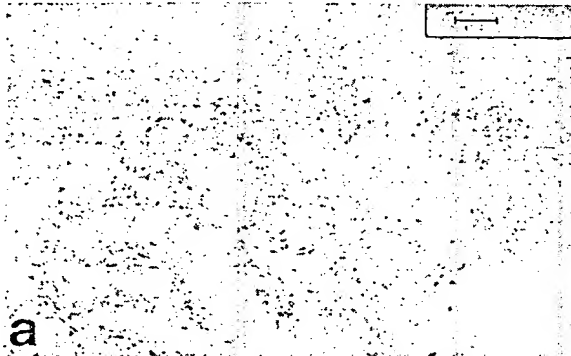


Figure 6

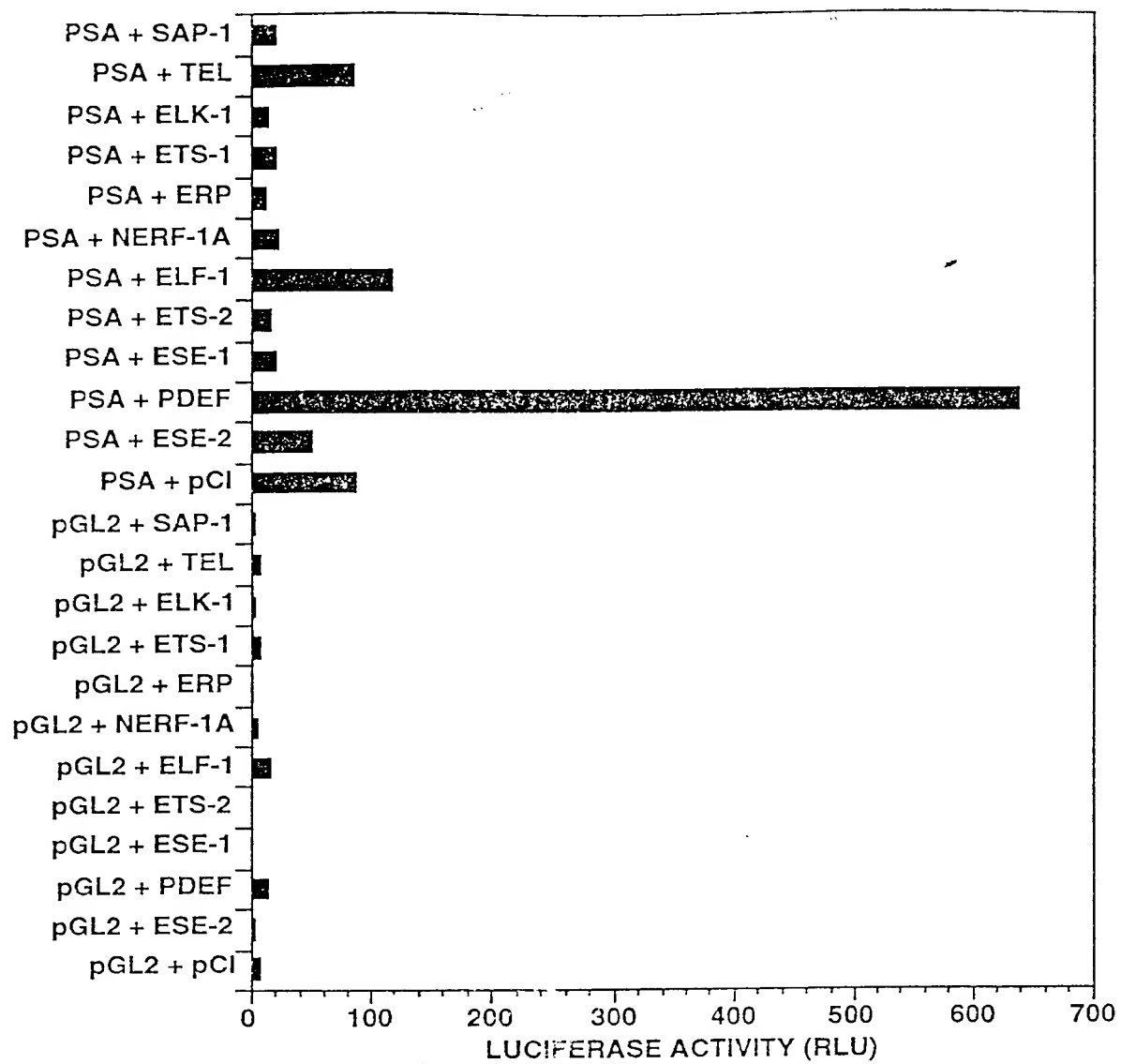


Figure 7

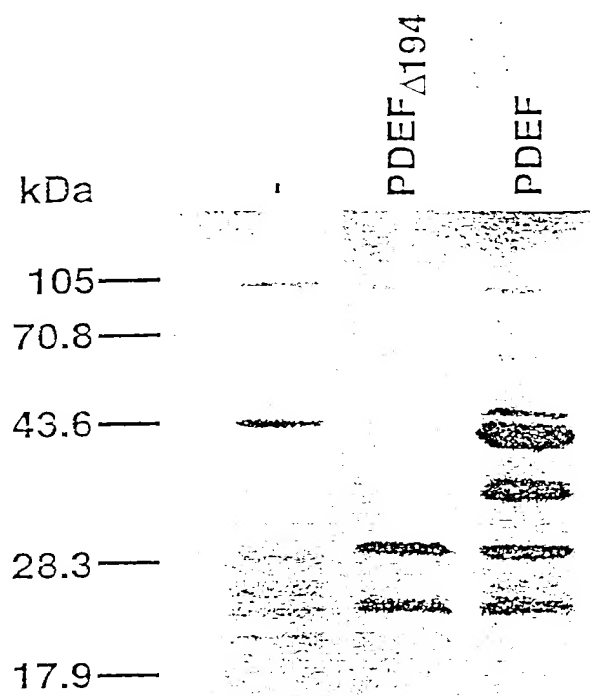


Figure 9

GST

GST-CTBP

GST-AR1-676

GST-AR505-676

GST-AR505-919

GST-AR6760919

GST-AR676-844

GST-AR875-919

GST-AR505-559

GST-AR559-624

GST-AR624-676

← PDEF

FIGURE 10

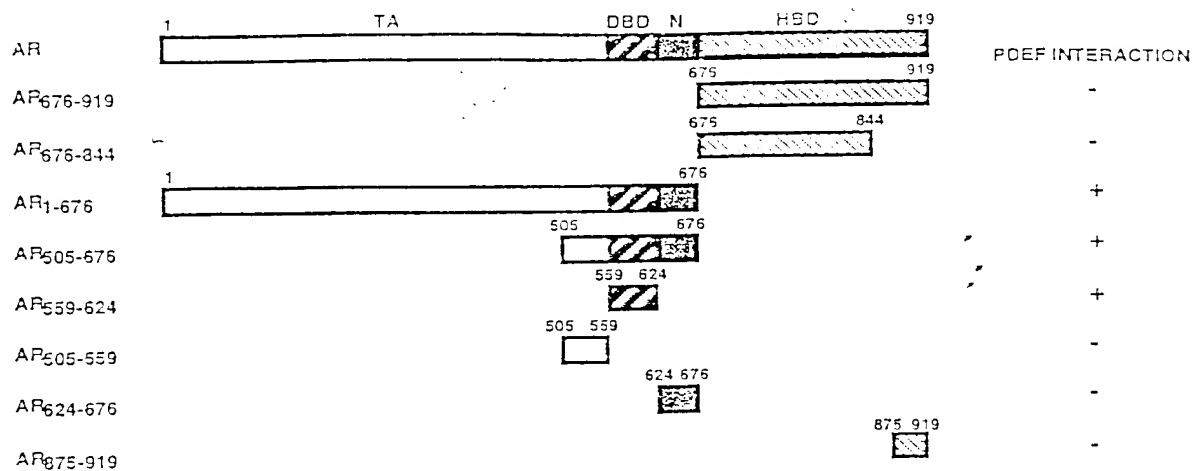


Figure 8

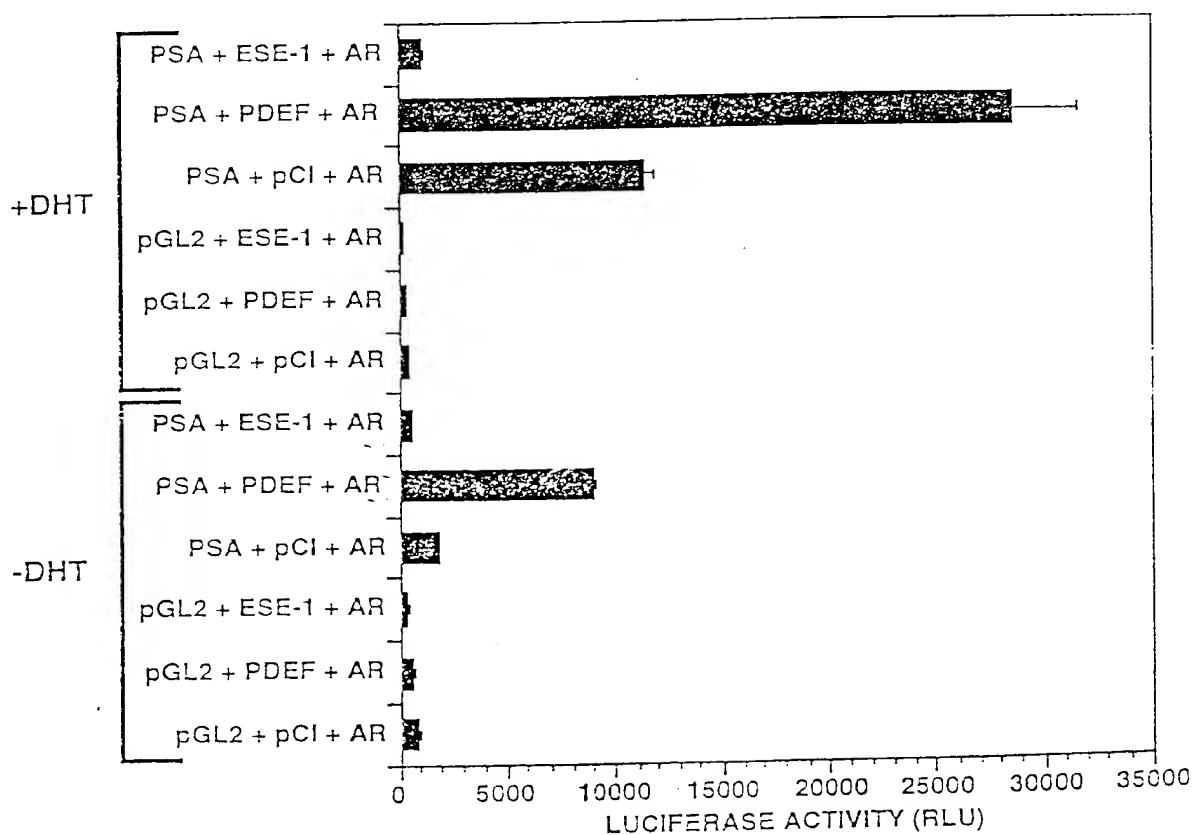


Figure 11